

chain nodes :

38

ring nodes :

1 2 3 4 5 6 7 8 9

ring/chain nodes :

10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30  
31 32 33 34 35 36 37

ring/chain bonds :

7-11 8-12 9-10 12-13 13-14 13-15 13-16 14-17 15-24 16-31 17-18 18-19  
19-20 20-21 21-22 22-23 24-25 25-26 26-27 27-28 28-29 29-30 31-32 32-33  
33-34 34-35 35-36 36-37

ring bonds :

1-2 1-6 2-3 3-4 4-5 4-7 5-6 5-9 7-8 8-9

exact/norm bonds :

12-13 13-14 13-15 13-16 14-17 15-24 16-31 17-18 18-19 19-20 20-21 21-22  
 22-23 24-25 25-26 26-27 27-28 28-29 29-30 31-32 32-33 33-34 34-35 35-36  
 36-37

exact bonds :

4-7 5-9 7-8 7-11 8-9 8-12 9-10

normalized bonds :

1-2 1-6 2-3 3-4 4-5 5-6

Match level :

1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:Atom 8:Atom 9:Atom 10:CLASS  
 11:CLASS 12:CLASS 13:CLASS 14:CLASS 15:CLASS 16:CLASS 17:CLASS 18:CLASS  
 19:CLASS 20:CLASS 21:CLASS 22:CLASS 23:CLASS 24:CLASS 25:CLASS 26:CLASS  
 27:CLASS 28:CLASS 29:CLASS 30:CLASS 31:CLASS 32:CLASS 33:CLASS 34:CLASS  
 35:CLASS 36:CLASS 37:CLASS 38:CLASS

L1 STRUCTURE UPLOADED

=> s l1 sss full

FULL SEARCH INITIATED 11:25:59 FILE 'REGISTRY'

FULL SCREEN SEARCH COMPLETED - 6 TO ITERATE

100.0% PROCESSED 6 ITERATIONS

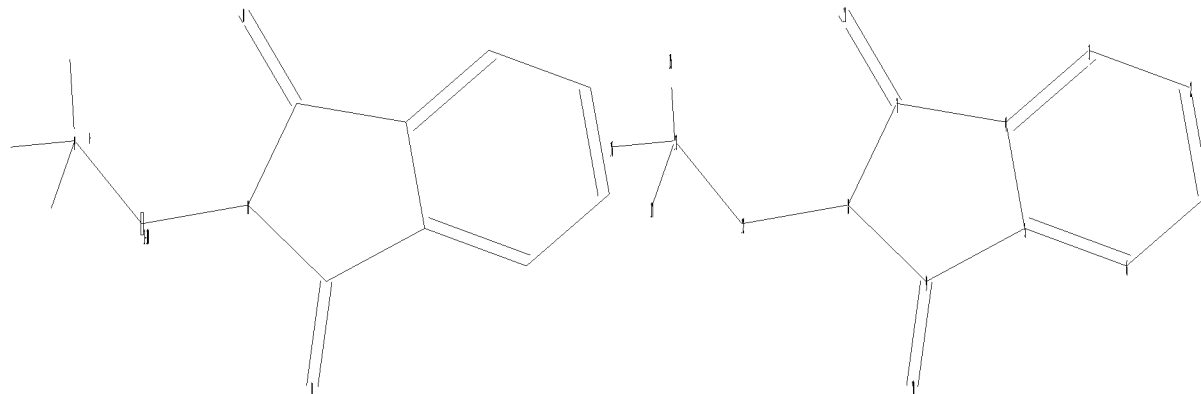
0 ANSWERS

SEARCH TIME: 00.00.01

L2 0 SEA SSS FUL L1

=>

Uploading C:\Documents and Settings\mpepitone\My Documents\ChemDraw\11663870\2.str



chain nodes :

15 16 17

ring nodes :

1 2 3 4 5 6 7 8 9

ring/chain nodes :

10 11 12 13

chain bonds :

13-15 13-16 13-17

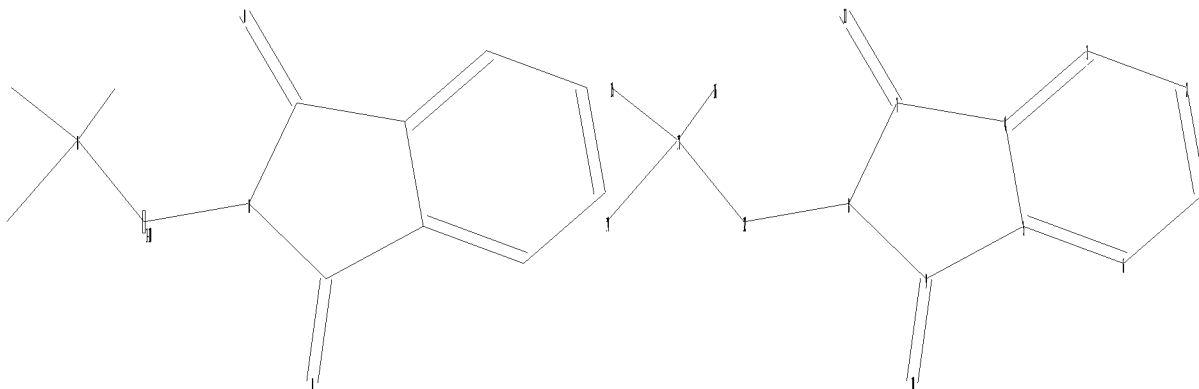
ring/chain bonds :

7-11 8-12 9-10 12-13

ring bonds :

1-2 1-6 2-3 3-4 4-5 4-7 5-6 5-9 7-8 8-9

exact/norm bonds :



```

chain nodes :
15 16
ring nodes :
1 2 3 4 5 6 7 8 9
ring/chain nodes :
10 11 12 13 14
chain bonds :
13-15 13-16
ring/chain bonds :
7-11 8-12 9-10 12-13 13-14
ring bonds :
1-2 1-6 2-3 3-4 4-5 4-7 5-6 5-9 7-8 8-9
exact/norm bonds :
12-13 13-14 13-15 13-16
exact bonds :
4-7 5-9 7-8 7-11 8-9 8-12 9-10
normalized bonds :
1-2 1-6 2-3 3-4 4-5 5-6

```

```

Match level :
1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:Atom 8:Atom 9:Atom 10:CLASS
11:CLASS 12:CLASS 13:CLASS 14:CLASS 15:CLASS 16:CLASS

```

L6 STRUCTURE UPLOADED

```

=> s 16 sss full
FULL SEARCH INITIATED 11:38:20 FILE 'REGISTRY'
FULL SCREEN SEARCH COMPLETED - 3032 TO ITERATE

```

```

100.0% PROCESSED      3032 ITERATIONS      572 ANSWERS
SEARCH TIME: 00.00.01

```

L7 572 SEA SSS FUL L6

```

=> file caplus
COST IN U.S. DOLLARS      SINCE FILE      TOTAL
                           ENTRY      SESSION
FULL ESTIMATED COST      191.54      654.53

DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS)      SINCE FILE      TOTAL

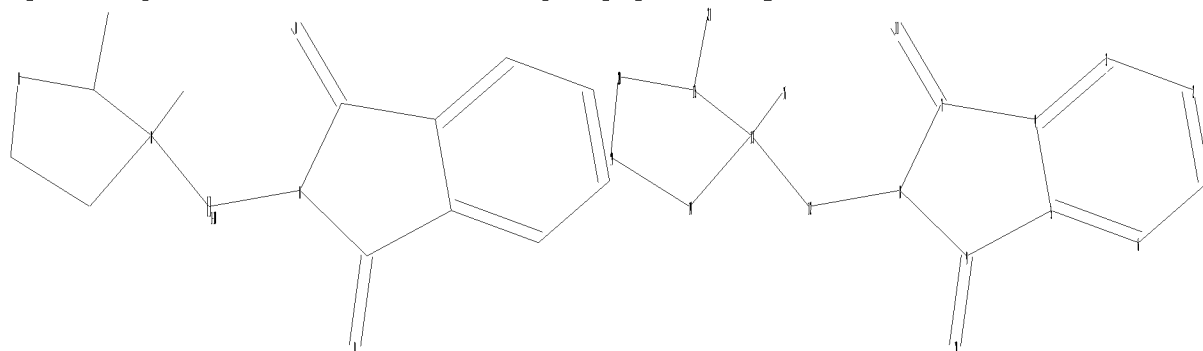
```

on property searching in REGISTRY, refer to:

<http://www.cas.org/support/stngen/stdoc/properties.html>

=>

Uploading C:\Documents and Settings\mpepitone\My Documents\ChemDraw\11663870\6.str



chain nodes :

16 21

ring nodes :

1 2 3 4 5 6 7 8 9 13 14 15 19 20

ring/chain nodes :

10 11 12

chain bonds :

13-16 15-21

ring/chain bonds :

7-11 8-12 9-10 12-13

ring bonds :

1-2 1-6 2-3 3-4 4-5 4-7 5-6 5-9 7-8 8-9 13-14 13-15 14-19 15-20 19-20

exact/norm bonds :

12-13 13-14 13-15 13-16 14-19 15-20 19-20

exact bonds :

4-7 5-9 7-8 7-11 8-9 8-12 9-10 15-21

normalized bonds :

1-2 1-6 2-3 3-4 4-5 5-6

Match level :

1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:Atom 8:Atom 9:Atom 10:CLASS

11:CLASS 12:CLASS 13:CLASS 14:CLASS 15:CLASS 16:CLASS 19:Atom 20:Atom

21:CLASS

L13 STRUCTURE UPLOADED

=> s l13 sss full

FULL SEARCH INITIATED 12:03:15 FILE 'REGISTRY'

FULL SCREEN SEARCH COMPLETED - 2 TO ITERATE

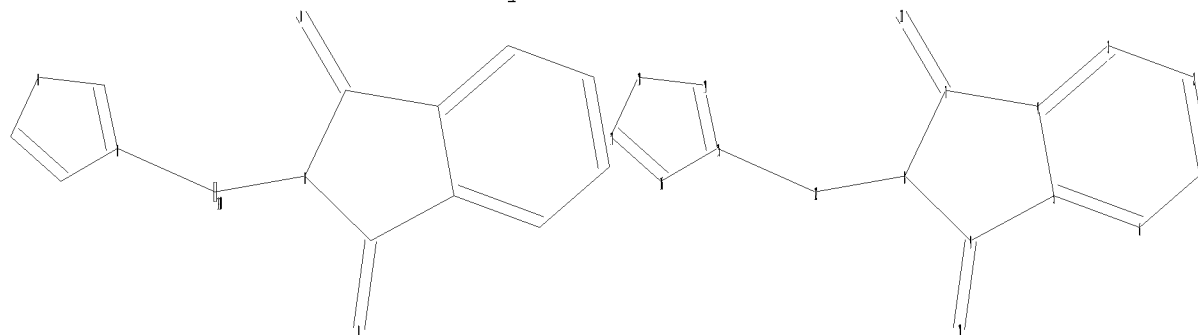
100.0% PROCESSED 2 ITERATIONS

0 ANSWERS

SEARCH TIME: 00.00.01

L14 0 SEA SSS FUL L13

=>  
 Uploading C:\Documents and Settings\mpepitone\My  
 Documents\ChemDraw\11663870\imuy.str



ring nodes :  
 1 2 3 4 5 6 7 8 9 15 16 17 18 19  
 ring/chain nodes :  
 10 11 12  
 chain bonds :  
 12-15  
 ring/chain bonds :  
 7-11 8-12 9-10  
 ring bonds :  
 1-2 1-6 2-3 3-4 4-5 4-7 5-6 5-9 7-8 8-9 15-16 15-19 16-17 17-18 18-19  
 exact/norm bonds :  
 12-15 15-16 15-19 16-17 17-18 18-19  
 exact bonds :  
 4-7 5-9 7-8 7-11 8-9 8-12 9-10  
 normalized bonds :  
 1-2 1-6 2-3 3-4 4-5 5-6

Match level :  
 1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:Atom 8:Atom 9:Atom 10:CLASS  
 11:CLASS 12:CLASS 15:Atom 16:Atom 17:Atom 18:Atom 19:Atom

L15 STRUCTURE UPLOADED

=> s l15 sss full  
 FULL SEARCH INITIATED 12:05:35 FILE 'REGISTRY'  
 FULL SCREEN SEARCH COMPLETED - 6915 TO ITERATE

100.0% PROCESSED 6915 ITERATIONS 57 ANSWERS  
 SEARCH TIME: 00.00.01

L16 57 SEA SSS FUL L15

=> file caplus		
COST IN U.S. DOLLARS	SINCE FILE	TOTAL
	ENTRY	SESSION
FULL ESTIMATED COST	384.55	2084.38